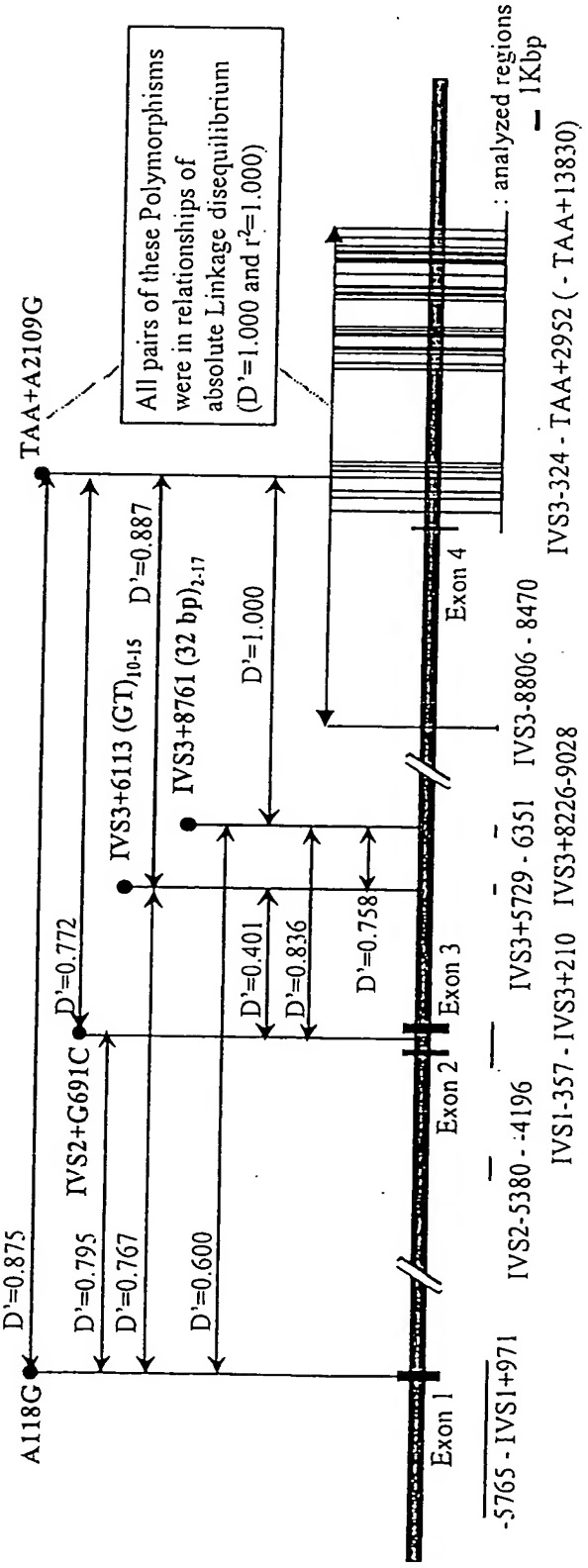
FIG. 1


All pairs of these Polymorphisms were in relationships of absolute Linkage disequilibrium (D'=1.000 and r²=1.000)
: analyzed regions
1Kbp
TAA+A2109G
A118G
IVS2+G691C
IVS3+6113 (GT)10-15
IVS3+8761 (32 bp)2-17
D'=0.875
D'=0.795
D'=0.767
D'=0.600
D'=0.772
D'=0.887
D'=1.000
D'=0.401
D'=0.836
D'=0.758
Exon 1
Exon 2
Exon 3
Exon 4
-5765 - IVS1+971
IVS2-5380 - -4196
IVS1-357 - IVS3+210
IVS3+5729 - 6351
IVS3+8226-9028
IVS3-8806 - 8470
IVS3-324 - TAA+2952 (- TAA+13830)